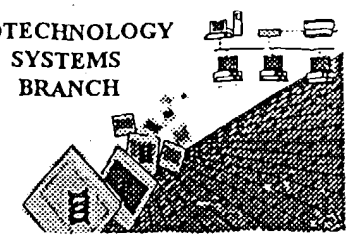
BIOTECHNOLOGY
SYSTEMS
BRANCH



# RAW SEQUENCE LISTING ERROR REPORT

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:**

Application Serial Number: 10/656,093B
Source: 1FWO
Date Processed by STIC: 4/14/04

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**
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*Effective 12/13/03*: **TELEPHONE: 571-272-2510; FAX: 571-273-0221**

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**http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm**

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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<**http://www.uspto.gov/ebc/efs/downloads/documents.htm**> , **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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Revised 10/08/03

IFWO

**RAW SEQUENCE LISTING** DATE: 04/14/2004
PATENT APPLICATION: **US/10/656,093B** TIME: 14:39:47

Input Set : **A:\256-152div corrected in response to notice to comply.txt**
Output Set: **N:\CRF4\04142004\J656093B.raw**

pp 4,6

<110> APPLICANT: YOUNG, ANDREW A.
VINE, WILL
BEELEY, NIGEL R.A.
PRICKETT, KATHRYN S.
<120> TITLE OF INVENTION: INOTROPIC AND DIURETIC EFFECTS OF GLP-1 AND GLP-1 AGONISTS
<130> FILE REFERENCE: 256-152DIV US
<140> CURRENT APPLICATION NUMBER: 10/656,093B
<141> CURRENT FILING DATE: 2003-09-05
<160> NUMBER OF SEQ ID NOS: 75
<170> SOFTWARE: PatentIn Ver. 2.1
<210> SEQ ID NO: 1
<211> LENGTH: 39
<212> TYPE: PRT
<213> ORGANISM: Heloderma horridum
<220> FEATURE:
<223> OTHER INFORMATION: Exendin-3
<400> SEQUENCE: 1

```
His Ser Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1               5                  10                  15
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
            20                  25                  30
Ser Gly Ala Pro Pro Pro Ser
        35
```

<210> SEQ ID NO: 2
<211> LENGTH: 39
<212> TYPE: PRT
<213> ORGANISM: Heloderma suspectum
<220> FEATURE:
<223> OTHER INFORMATION: Exendin-4
<400> SEQUENCE: 2

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His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 1               5                  10                  15
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
            20                  25                  30
Ser Gly Ala Pro Pro Pro Ser
        35
```

<210> SEQ ID NO: 3
<211> LENGTH: 30
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<220> FEATURE:
<223> OTHER INFORMATION: GLP-1
<400> SEQUENCE: 3

**RAW SEQUENCE LISTING** DATE: 04/14/2004
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```
66 His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly
67  1               5                  10                  15
69 Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg
70             20                  25                  30
73 <210> SEQ ID NO: 4
74 <211> LENGTH: 39
75 <212> TYPE: PRT
76 <213> ORGANISM: Artificial Sequence
78 <220> FEATURE:
79 <223> OTHER INFORMATION: Description of Artificial Sequence: Exendin or
80       exendin agonist
82 <220> FEATURE:
83 <221> NAME/KEY: MOD_RES
84 <222> LOCATION: (1)
85 <223> OTHER INFORMATION: His, Arg or Tyr
87 <220> FEATURE:
88 <221> NAME/KEY: MOD_RES
89 <222> LOCATION: (2)
90 <223> OTHER INFORMATION: Ser, Gly, Ala or Thr
92 <220> FEATURE:
93 <221> NAME/KEY: MOD_RES
94 <222> LOCATION: (3)
95 <223> OTHER INFORMATION: Asp or Glu
97 <220> FEATURE:
98 <221> NAME/KEY: MOD_RES
99 <222> LOCATION: (5)
100 <223> OTHER INFORMATION: Ala or Thr
102 <220> FEATURE:
103 <221> NAME/KEY: MOD_RES
104 <222> LOCATION: (6)
105 <223> OTHER INFORMATION: Ala, Phe, Tyr or naphthylalanine
107 <220> FEATURE:
108 <221> NAME/KEY: MOD_RES
109 <222> LOCATION: (7)
110 <223> OTHER INFORMATION: Thr or Ser
112 <220> FEATURE:
113 <221> NAME/KEY: MOD_RES
114 <222> LOCATION: (8)
115 <223> OTHER INFORMATION: Ala, Ser or Thr
117 <220> FEATURE:
118 <221> NAME/KEY: MOD_RES
119 <222> LOCATION: (9)
120 <223> OTHER INFORMATION: Asp or Glu
122 <220> FEATURE:
123 <221> NAME/KEY: MOD_RES
124 <222> LOCATION: (10)
125 <223> OTHER INFORMATION: Ala, Leu, Ile, Val, pentylglycine or Met
127 <220> FEATURE:
128 <221> NAME/KEY: MOD_RES
```

```
129 <222> LOCATION: (11)
130 <223> OTHER INFORMATION: Ala or Ser
132 <220> FEATURE:
133 <221> NAME/KEY: MOD_RES
134 <222> LOCATION: (12)
135 <223> OTHER INFORMATION: Ala or Lys
137 <220> FEATURE:
138 <221> NAME/KEY: MOD_RES
139 <222> LOCATION: (13)
140 <223> OTHER INFORMATION: Ala or Gln
142 <220> FEATURE:
143 <221> NAME/KEY: MOD_RES
144 <222> LOCATION: (14)
145 <223> OTHER INFORMATION: Ala, Leu, Ile, pentylglycine, Val or Met
147 <220> FEATURE:
148 <221> NAME/KEY: MOD_RES
149 <222> LOCATION: (15)...(17)
150 <223> OTHER INFORMATION: Ala or Glu
152 <220> FEATURE:
153 <221> NAME/KEY: MOD_RES
154 <222> LOCATION: (19)
155 <223> OTHER INFORMATION: Ala or Val
157 <220> FEATURE:
158 <221> NAME/KEY: MOD_RES
159 <222> LOCATION: (20)
160 <223> OTHER INFORMATION: Ala or Arg
162 <220> FEATURE:
163 <221> NAME/KEY: MOD_RES
164 <222> LOCATION: (21)
165 <223> OTHER INFORMATION: Ala or Leu
167 <220> FEATURE:
168 <221> NAME/KEY: MOD_RES
169 <222> LOCATION: (22)
170 <223> OTHER INFORMATION: Phe, Tyr or naphthylalanine
172 <220> FEATURE:
173 <221> NAME/KEY: MOD_RES
174 <222> LOCATION: (23)
175 <223> OTHER INFORMATION: Ile, Val, Leu, pentylglycine, tert-butylglycine or Met
177 <220> FEATURE:
178 <221> NAME/KEY: MOD_RES
179 <222> LOCATION: (24)
180 <223> OTHER INFORMATION: Ala, Glu or Asp
182 <220> FEATURE:
183 <221> NAME/KEY: MOD_RES
184 <222> LOCATION: (25)
185 <223> OTHER INFORMATION: Ala, Trp, Phe, Tyr or naphthylalanine
187 <220> FEATURE:
188 <221> NAME/KEY: MOD_RES
189 <222> LOCATION: (26)
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**RAW SEQUENCE LISTING** DATE: 04/14/2004
PATENT APPLICATION: **US/10/656,093B** TIME: 14:39:47

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Output Set: **N:\CRF4\04142004\J656093B.raw**

what about Xaa's at locations 29-30, 32-35? (see below)

```
190 <223> OTHER INFORMATION: Ala or Leu
192 <220> FEATURE:
193 <221> NAME/KEY: MOD_RES
194 <222> LOCATION: (27)
195 <223> OTHER INFORMATION: Ala or Lys
197 <220> FEATURE:
198 <221> NAME/KEY: MOD_RES
199 <222> LOCATION: (28)
200 <223> OTHER INFORMATION: Ala or Asn
202 <220> FEATURE:
203 <221> NAME/KEY: MOD_RES
204 <222> LOCATION: (31)
205 <223> OTHER INFORMATION: Pro, homoproline, 3Hyp, 4Hyp, thioproline, N-alkylglycine
206       N-alkylpentylglycine or N-alkylalanine
208 <220> FEATURE:
209 <221> NAME/KEY: MOD_RES
210 <222> LOCATION: (36)..(38)
211 <223> OTHER INFORMATION: Pro, homoproline, 3Hyp, 4Hyp, thioproline, N-alkylglycine
212       N-alkylpentylglycine or N-alkylalanine
214 <220> FEATURE:
215 <221> NAME/KEY: MOD_RES
216 <222> LOCATION: (39)
217 <223> OTHER INFORMATION: Ser, Thr, Tyr, Pro, homoproline, 3Hyp, 4Hyp, thioproline,
218       N-alkylglycine, N-alkylpentylglycine or N-alkylalanine
220 <220> FEATURE:
221 <223> OTHER INFORMATION: provided no more than three of Xaa5, Xaa6, Xaa8,
222       Xaa10, Xaa11, Xaa12, Xaa13, Xaa14, Xaa15, Xaa16, Xaa17,
223       Xaa19, Xaa20, Xaa21, Xaa24, Xaa25, Xaa26, Xaa27 or Xaa28
224       are Ala; and the compound is not exendin-3 or exendin-4
226 <220> FEATURE:
227 <223> OTHER INFORMATION: this peptide may encompass 28-39 residues, wherein
228       residues 1-28 are constant and residues 29-39 may vary
229       in length according to the specification
231 <400> SEQUENCE: 4
W--> 232 Xaa Xaa Xaa Gly Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
233   1               5                  10                  15
235 Xaa Ala Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
236             20                  25                  30
238 Xaa Xaa Xaa Xaa Xaa Xaa Xaa
239         35
241 <210> SEQ ID NO: 5
242 <211> LENGTH: 30
243 <212> TYPE: PRT
244 <213> ORGANISM: Artificial Sequence
246 <220> FEATURE:
247 <223> OTHER INFORMATION: Description of Artificial Sequence: Exendin or
248       GLP-1 agonist
250 <220> FEATURE:
251 <223> OTHER INFORMATION: C-term may be amidated
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**RAW SEQUENCE LISTING** DATE: 04/14/2004
PATENT APPLICATION: **US/10/656,093B** TIME: 14:39:47

Input Set : **A:\256-152div corrected in response to notice to comply.txt**
Output Set: **N:\CRF4\04142004\J656093B.raw**

253 <400> SEQUENCE: 5
254 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
255 1 5 10 15
257 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly
258 20 25 30
261 <210> SEQ ID NO: 6
262 <211> LENGTH: 28
263 <212> TYPE: PRT
264 <213> ORGANISM: Artificial Sequence
266 <220> FEATURE:
267 <223> OTHER INFORMATION: Description of Artificial Sequence: Exendin or
268 GLP-1 agonist
270 <220> FEATURE:
271 <223> OTHER INFORMATION: C-term amidated
273 <400> SEQUENCE: 6
274 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
275 1 5 10 15
277 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
278 20 25
281 <210> SEQ ID NO: 7
282 <211> LENGTH: 28
283 <212> TYPE: PRT
284 <213> ORGANISM: Artificial Sequence
286 <220> FEATURE:
287 <223> OTHER INFORMATION: Description of Artificial Sequence: Exendin or
288 GLP-1 agonist
290 <220> FEATURE:
291 <223> OTHER INFORMATION: C-term amidated
293 <400> SEQUENCE: 7
294 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
295 1 5 10 15
297 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
298 20 25
301 <210> SEQ ID NO: 8
302 <211> LENGTH: 28
303 <212> TYPE: PRT
304 <213> ORGANISM: Artificial Sequence
306 <220> FEATURE:
307 <223> OTHER INFORMATION: Description of Artificial Sequence: Exendin or
308 GLP-1 agonist
310 <220> FEATURE:
311 <223> OTHER INFORMATION: C-term amidated
313 <400> SEQUENCE: 8
314 His Ala Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
315 1 5 10 15
317 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
318 20 25
321 <210> SEQ ID NO: 9
322 <211> LENGTH: 28

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 04/14/2004
PATENT APPLICATION: US/10/656,093B TIME: 14:39:48

Input Set : A:\256-152div corrected in response to notice to comply.txt
Output Set: N:\CRF4\04142004\J656093B.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:4; Xaa Pos. 1,2,3,5,6,7,8,9,10,11,12,13,14,15,16,17,18,20,21,22,23,24
Seq#:4; Xaa Pos. 25,26,27,28,29,30,31,32,33,34,35,36,37,38,39
Seq#:47; Xaa Pos. 31,36,37,38
Seq#:48; Xaa Pos. 36,37,38
Seq#:49; Xaa Pos. 31
Seq#:50; Xaa Pos. 31,36,37
Seq#:51; Xaa Pos. 31,36,37
Seq#:52; Xaa Pos. 31,36
Seq#:55; Xaa Pos. 6
Seq#:59; Xaa Pos. 10
Seq#:60; Xaa Pos. 22
Seq#:61; Xaa Pos. 23
Seq#:65; Xaa Pos. 31,36,37
Seq#:66; Xaa Pos. 19
Seq#:67; Xaa Pos. 17
Seq#:75; Xaa Pos. 29

**VERIFICATION SUMMARY** DATE: 04/14/2004
PATENT APPLICATION: **US/10/656,093B** TIME: 14:39:48

Input Set : **A:\256-152div corrected in response to notice to comply.txt**
Output Set: **N:\CRF4\04142004\J656093B.raw**

L:232 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
M:341 Repeated in SeqNo=4
L:1131 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:16
M:341 Repeated in SeqNo=47
L:1162 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:32
L:1187 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49 after pos.:16
L:1220 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 after pos.:16
M:341 Repeated in SeqNo=50
L:1253 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 after pos.:16
M:341 Repeated in SeqNo=51
L:1286 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52 after pos.:16
M:341 Repeated in SeqNo=52
L:1354 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:0
L:1439 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:0
L:1467 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60 after pos.:16
L:1492 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:16
L:1585 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65 after pos.:16
M:341 Repeated in SeqNo=65
L:1635 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66 after pos.:16
L:1642 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:67
L:1673 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:16
L:1878 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:75 after pos.:16